



HR - skin2.ST25
SEQUENCE LISTING

<110> The Procter & Gamble Company

<120> Hairless Protein-Interacting Partner Complexes And Methods
Thereof For The Beautification And/Or Improvement Of Mammalian
Skin

<130> 9423

<160> 16

<170> PatentIn version 3.2

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<212> DNA

<213> Homo Sapiens Keratin 5

<400> 1

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<213> Homo Sapiens Ubiquitous Receptor

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 <211> 792
 <212> DNA
 <213> Homo Sapiens Similar to Stromal Antigen 2

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 <212> DNA
 <213> Homo Sapiens Retinoic Acid Receptor Gamma-1

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 <213> Homo Sapiens Thyroid Hormone Receptor Alpha

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 <213> Homo Sapiens HIC Protein Isoform P32 and Isoform 40

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 <211> 610
 <212> DNA
 <213> Homo Sapiens Insulin-like Growth Factor Binding Domain Protein 6

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 <212> DNA
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 <212> DNA
 <213> Homo Sapiens Endoplasmic reticulum thioredoxin superfamily member

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<210> 13
 <211> 779
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 <213> Homo Sapiens Protein Inhibitor of Activated STAT-3

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<210> 14
 <211> 738
 <212> DNA
 <213> Homo Sapiens DEAD box polypeptide 3

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<210> 15
 <211> 450
 <212> DNA
 <213> Homo Sapiens Dpy-30 Like Protein

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 <211> 422
 <212> PRT
 <213> Mouse VDR

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 50 55 60
 Asp Cys Arg Ile Thr Lys Asp Asn Arg Arg His Cys Gln Ala Cys Arg
 65 70 75 80
 Leu Lys Arg Cys Val Asp Ile Gly Met Met Lys Glu Phe Ile Leu Thr
 85 90 95
 Asp Glu Glu Val Gln Arg Lys Arg Glu Met Ile Met Lys Arg Lys Glu
 100 105 110

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Glu Glu Ala Leu Lys Asp Ser Leu Arg Pro Lys Leu Ser Glu Glu Gln
115 120 125

Gln His Ile Ile Ala Ile Leu Leu Asp Ala His His Lys Thr Tyr Asp
130 135 140

Pro Thr Tyr Ala Asp Phe Arg Asp Phe Arg Pro Pro Ile Arg Ala Asp
145 150 155 160

Val Ser Thr Gly Ser Tyr Ser Pro Arg Pro Thr Leu Ser Phe Ser Gly
165 170 175

Asp Ser Ser Ser Asn Ser Asp Leu Tyr Thr Pro Ser Leu Asp Met Met
180 185 190

Glu Pro Ala Ser Phe Ser Thr Met Asp Leu Asn Glu Glu Gly Ser Asp
195 200 205

Asp Pro Ser Val Thr Leu Asp Leu Ser Pro Leu Ser Met Leu Pro His
210 215 220

Leu Ala Asp Leu Val Ser Tyr Ser Ile Gln Lys Val Ile Gly Phe Ala
225 230 235 240

Lys Met Ile Pro Gly Phe Arg Asp Leu Thr Ser Asp Asp Gln Ile Val
245 250 255

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260 265 270

Ser Phe Thr Leu Asp Asp Met Ser Trp Asp Cys Gly Ser Gln Asp Tyr
275 280 285

Lys Tyr Asp Ile Thr Asp Val Ser Arg Ala Gly His Thr Leu Glu Leu
290 295 300

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His Glu Glu Glu His Val Leu Leu Met Ala Ile Cys Ile Val Ser Pro
325 330 335

Asp Arg Pro Gly Val Gln Asp Ala Lys Leu Val Glu Ala Ile Gln Asp
340 345 350

Arg Leu Ser Asn Thr Leu Gln Thr Tyr Ile Arg Cys Arg His Pro Pro
355 360 365

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Pro Gly Ser His Gln Leu Tyr Ala Lys Met Ile Gln Lys Leu Ala Asp
 370 375 380

Leu Arg Ser Leu Asn Glu Glu His Ser Lys Gln Tyr Arg Ser Leu Ser
 385 390 395 400

Phe Gln Pro Glu Asn Ser Met Lys Leu Thr Pro Leu Val Leu Glu Val
 405 410 415

Phe Gly Asn Glu Ile Ser
 420